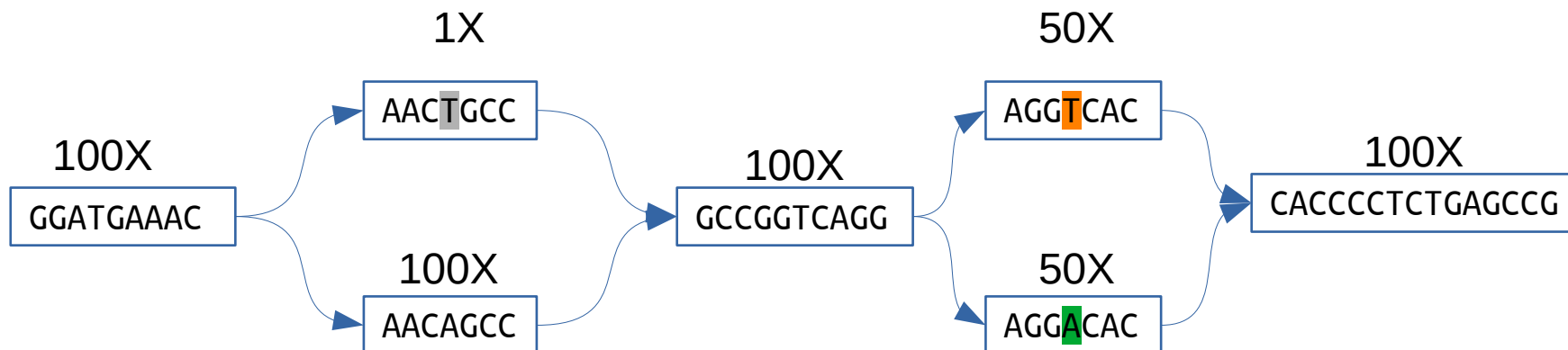


Sequencing error

Heterozygous variant



Quasi repeats

